

0500

OIPE

RAW SEQUENCE LISTING  
 PATENT APPLICATION: US/09/553,094

DATE: 05/09/2000  
 TIME: 16:28:06

Input Set : D:\AprilCornReg.rpt  
 Output Set: N:\CRF3\05092000\I553094.raw

ENTERED

1 <110> APPLICANT: Andersen, Scott E.  
 2 Fisher, Dane K.  
 4 <120> TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With  
 5 Plants  
 7 <130> FILE REFERENCE: 38-21(15502)B  
 C--> 9 <140> CURRENT APPLICATION NUMBER: US/09/553,094  
 C--> 9 <141> CURRENT FILING DATE: 2000-04-18  
 9 <160> NUMBER OF SEQ ID NOS: 18831  
 11 <210> SEQ ID NO: 1  
 12 <211> LENGTH: 341  
 13 <212> TYPE: DNA  
 14 <213> ORGANISM: Zea mays  
 16 <223> OTHER INFORMATION: Clone ID: LIB3061-001-Q1-K2-H1  
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 22 cggaaaggatc ggctggctcg tggtcagggt cgctctgcag agcgaggatg tcgagctcgt 120  
 24 ctgctcaac gaccgcttca tcaccacgga ttacatgacc tacatgttca agtacgacac 180  
 26 cgtgcacggt caatggaagc acatgcgaca ttaccctcaa ggactccaag acgcttcttg 240  
 28 ttggcgagaa gccggtcacc gtctttggca tcaagaacct ccgagagatt ccgtgatgtg 300  
 30 aggctggtgc tgtatatgtc gtgcagtcga tcgggtgtctt c 341  
 33 <210> SEQ ID NO: 2  
 34 <211> LENGTH: 371  
 35 <212> TYPE: DNA  
 36 <213> ORGANISM: Zea mays  
 38 <223> OTHER INFORMATION: Clone ID: LIB3061-005-Q1-K1-B3  
 40 <400> SEQUENCE: 2  
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 44 tcaccctcat ggccaagatc aagatcgga tcaatgggtt tggcaggatc ggaaggctgg 120  
 46 tggccagggt ggccttcgag agcgacgacg tcgagctcgt cggcgtgaac gacccttca 180  
 48 tcagcacgga ttacatgacc tacatgttca agtatgacac tgtgcacgga cagtgggaagc 240  
 50 accatgaggt caaggtgaag gactccaaga cccttctctt tgggtgagaa gaggttgctg 300  
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 54 ttgagtctac t 371  
 57 <210> SEQ ID NO: 3  
 58 <211> LENGTH: 328  
 59 <212> TYPE: DNA  
 60 <213> ORGANISM: Zea mays  
 62 <223> OTHER INFORMATION: Clone ID: LIB3061-056-Q1-K1-F11  
 64 <400> SEQUENCE: 3  
 66 acgcccgtcat cgatctcgtt tcaaatccat ccacccctcg ttccctccgc ccgcagcacc 60  
 68 tcaaccctgat ggggaagatc aagatcgga tcaatgggtt cggcaggatc ggaaggctgg 120  
 70 tggccagggt ggccttcgag agtgacgatg tcgagctcgt cggcgtgaac gacccttca 180  
 72 tcagcacgga ctacatgacc tacatgttca agtacgacac tgtgcacgga cagtgggaagc 240  
 74 accatgaagt caaggtgaag gactccaaaa cccttctctt tgggtgagaag gaagttgctg 300  
 76 tgtttggtcg caggaaccct gaggagat 328  
 79 <210> SEQ ID NO: 4  
 80 <211> LENGTH: 408

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81 <212> TYPE: DNA
82 <213> ORGANISM: Zea mays
84 <223> OTHER INFORMATION: Clone ID: LIB3061-013-Q1-K1-D10
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90 tgcctccgt ctacgagcct ctcaccctca tggccaagat caagatcggg atcaatgggt 120
92 tcggcaggat cgggaaggctg gtggccaggg tggccctgca gagcgacgac gtcgagctcg 180
94 tcgccgtgaa cgacccttc atcagcaccg attacatgac ctacatgttc aagtatgaca 240
96 ctgtgcacgg acagtggagg caccatgagg tcaagggtgaa ggactccaag acccttctct 300
98 ttggtgagaa ggaggtcgct gtgtttggct gcaggaaccc tgaggagatc ccatggtgct 360
100 cggttggcgc tgaatatgct gtcgagtcta ctggtgtttt cactgacc 408
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104 <211> LENGTH: 417
105 <212> TYPE: DNA
106 <213> ORGANISM: Zea mays
108 <223> OTHER INFORMATION: Clone ID: LIB3061-011-Q1-K1-H1
110 <400> SEQUENCE: 5
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114 tagcagcctc tcaccctcat ggccaagatc aagatcggga tcaatgggtt tggcaggatc 120
116 ggaaggcttg tggccagggt ggccctgcag agcgacgacg tcgagctcgt cgccgtgaac 180
118 gacccttca tcagcaccga ttacatgacc tacatgttca agtatgacac tgtgcacgga 240
120 cagtgggaag accatgagggt caagggtgaa gactccaaga cccttctctt tggtgagaag 300
122 gaagtgtgct gtgtttgctg caggaaccct gaggagatcc catgggggtc ggttggcgct 360
124 gaatatgttg ctgagctcac tgggtgtttc actgaccagg agaaggctgc agctcac 417
127 <210> SEQ ID NO: 6
128 <211> LENGTH: 422
129 <212> TYPE: DNA
130 <213> ORGANISM: Zea mays
132 <223> OTHER INFORMATION: Clone ID: LIB3061-034-Q1-K1-H6
134 <400> SEQUENCE: 6
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138 ctacgagcct ctcaccctca tggccaagat caagatcggg atcaatgggt tggcaggat 120
140 cgggaaggctg gtggccaggg tggccctgca gagcgacgac gtcgagctcg tcgccgtgaa 180
142 cgacccttc atcagcaccg attacatgac ctacatgttc aagtatgaca ctgtgcacgg 240
144 acagtggagg caccatgagg tcaagggtgaa ggactccaag acccttctct ttggtgagaa 300
146 ggaggttgct gtgtttggct gcaggaaccc tgaggagatc ccatgggggt cggttggcgc 360
148 tgaatatgtt gttgagtcta ctggtgtttt cactgaccag gagaaggctg cagctcactt 420
150 ga 422
153 <210> SEQ ID NO: 7
154 <211> LENGTH: 420
155 <212> TYPE: DNA
156 <213> ORGANISM: Zea mays
158 <223> OTHER INFORMATION: Clone ID: LIB3061-011-Q1-K1-D5
160 <400> SEQUENCE: 7
162 tcggaatcaa cggcttcgga aggatcgcca ggctcgtggc caggggtgcc ctccagagcg 60
164 aggatgtcga gtcgctgcc gtcaacgacc cttcatcac cacggactac atgacctaca 120
166 tgttcaagta cgacaccgtc cacggccact ggaagcacag cgacatcaca ctcaaggact 180
168 ccaagaccct tctcttcggt gacaagccgg tcaccgtctt tggcatcagg aacctgagg 240
170 aaatcccggt gggtaggct ggcgtgagt atgctgtgga gtccaccgac gtcttcactg 300

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172 acaaggacaa ggctgctgca catctcaagg gtggtgccaa gcaagctgat atctctgccc 360
174 caagcaaaaga cgcacccatg ttgacagttg gtgtcaatga ggacaagtac acctcggatg 420
177 <210> SEQ ID NO: 8
178 <211> LENGTH: 403
179 <212> TYPE: DNA
180 <213> ORGANISM: Zea mays
182 <223> OTHER INFORMATION: unsure at all n locations
183 <223> OTHER INFORMATION: Clone ID: LIB3061-034-Q1-K1-A7
185 <400> SEQUENCE: 8
187 ctcatcttgaa atcaacccat cctcgtttcc tccgtctagc agcctctcac ccgcatggcc 60
189 aagatcaaaga tcgggatcaa tgggtttggc aggatcggaa ggctggtggc cagggtggcc 120
191 ctgcagagcgc acgacgtcga gctcgtcgcc gtgaacgacc ccttcacag caccgattac 180
193 atgacctaca tgttcaagta tgacactgtg caccggacagt ggaagcacca tgaagtcaag 240
195 gtgaaggact ccaagaccct tctctttggt gagaaggagg ttgctgtgtt tggctgcagg 300
197 aacctgagg agatcccatg gggctcggtt ggcgtgaat atgttgctga gtctactggt 360
W--> 199 gttttcactg accaggagaa ggctgcagct cacttgaang gtg 403
202 <210> SEQ ID NO: 9
203 <211> LENGTH: 334
204 <212> TYPE: DNA
205 <213> ORGANISM: Zea mays
207 <223> OTHER INFORMATION: unsure at all n locations
208 <223> OTHER INFORMATION: Clone ID: LIB3061-037-Q1-K1-C8
210 <400> SEQUENCE: 9
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214 ggaggaatag tccactggag gtacatgagc ctaaaataat ttgtggagtg gtggttgact 120
216 tcagtggtaa gtgcctttca gactgtaccg gcatgcccaa aactgaaatt tccggtttag 180
218 atagaattgg gggcgcgga gaaggtgcca tgtgtgtgtt tgtgtgtgtt gcctgtggat 240
220 ggaggaattg atgggagtat gtgaactgat gtagctagtc tatgtttgtt accattgagg 300
222 gcttgttcgg ttagctttca atccatgtgg attg 334
225 <210> SEQ ID NO: 10
226 <211> LENGTH: 370
227 <212> TYPE: DNA
228 <213> ORGANISM: Zea mays
230 <223> OTHER INFORMATION: Clone ID: LIB3061-049-Q1-K1-G5
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234 atcccgtcgc atcccatccc agtaccaccc atctcgtcgt ccgccgattc gatccggttc 60
236 ttccaaccca ggggctgtgg gggcgacgaa gaggagcgtg ggcaccctcg gcgagcgga 120
238 tctgaagggtg aagaagggtg tctgtcgcgc cgacctcaac gtccctctcg acgacgcccc 180
240 gaagatcacc gacgacaccc gcatccgcgc ctacgtcccc accatcaagt tctcctcga 240
242 gaaggggcgc aaggctcatc tcggcagcca cctgtgtcgt gcaaaagggtg tcaccctaaa 300
244 gtacagcttg aagcctcttg atccacgctt gtctgagctc cttggagttg aagtcgtgat 360
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249 <210> SEQ ID NO: 11
250 <211> LENGTH: 435
251 <212> TYPE: DNA
252 <213> ORGANISM: Zea mays
254 <223> OTHER INFORMATION: Clone ID: LIB3061-006-Q1-K1-B5
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258 ccttcaagct ggtcaccaag cccgcggcgg cgctgcccgc tgctgctcac tggggagagc 60

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260 tcgcccgggg gccgcagggt accagccgcg ttggctcttg acacaagccc acaaacacag 120
262 ggcgctccgc acggacaaaa tctccgaaac gcccatggct aaggaagaga gggttgatcg 180
264 aagtgaataa ttgacatggg atagcatcag acaagttttg attagactag aagacagcat 240
266 catatttggg cttttggaga gagcacagtt ttgttacaac gctgacacat atgatagcaa 300
268 tgctttccac atggatgggt ttggagggtc tttggttgaa tatatggtta gagaaactga 360
270 aaagctccat gcacaggttg ggaggtacaa gagcccagat gagcaccctt tcttttccaa 420
272 ggatctgcct gagcc
275 <210> SEQ ID NO: 12
276 <211> LENGTH: 379
277 <212> TYPE: DNA
278 <213> ORGANISM: Zea mays
280 <223> OTHER INFORMATION: Clone ID: LIB3061-035-Q1-K1-B12
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286 atggccttca agctggtcac caagcccgcg gcgcgctcgc ccgctgctgc tcaactggga 120
288 gatctcgccc ggtggccgca ggttacgagc cgcgttgctc tcggaccagc gcccaggaa 180
290 aaggggctcc gcacgggcaa caactccgca acgcccattg ctaaggaaga gagggttgat 240
292 cgaagtgaaa tattgacatt ggtatgcatc agacaagttt tgattagact agaagacagc 300
294 atcatatttg gacttttggg gagagcacag ttttggtaca acgctgacac atatgatagc 360
296 aatgctatcc acatggatg
299 <210> SEQ ID NO: 13
300 <211> LENGTH: 392
301 <212> TYPE: DNA
302 <213> ORGANISM: Zea mays
304 <223> OTHER INFORMATION: Clone ID: LIB3061-034-Q1-K1-D8
306 <400> SEQUENCE: 13
308 caaattcata ctttcaatca gagccagtat cctcgcattg ttaccgagga cttcttgcca 60
310 cttcccagca aagggaaatc tgggaaggat ggctggtatc ctcttggtca tggatgatgtg 120
312 ttccctctt tgaataacag cggaaaactt gacatcttat tggctcaggg caaagagtat 180
314 gtctttgttg caaactcaga caacttgggt gctatagtcg acatcaagat cctaaacct 240
316 ctgatcaata accagaacga gtactgcatg gaagtactc caaagacgct ggctgacgtt 300
318 aaagtgggca ctctcatctc ttacgaaagg aagagttcag ctttttgagg atgcccagg 360
320 taccgatga gcatgtgaat gaatttaaat ca
323 <210> SEQ ID NO: 14
324 <211> LENGTH: 369
325 <212> TYPE: DNA
326 <213> ORGANISM: Zea mays
328 <223> OTHER INFORMATION: unsure at all n locations
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331 <400> SEQUENCE: 14
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335 gtccaccttg aaatgactgg gcagaacgtg accgagtgcg tcggtggatc acggaccgtg 120
337 accttcgacg atctgagcga ccgctaccac acccactcgc acccaaggct gaatgcctcc 180
339 cagtcctcgg agctcgcctt tatcatcgca gagaggctga ggaagaggag gatgcgatcg 240
W--> 341 gggtcaaca gcagctgcc actgcgcnca ctggctttct gagtagccgg agccaaacac 300
343 aaagggagggt aggaatagct gtggtgacct cggaggagaa gagacagtcg acgccttggt 360
345 tggatgatgc
348 <210> SEQ ID NO: 15
349 <211> LENGTH: 362

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350 <212> TYPE: DNA
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353 <223> OTHER INFORMATION: Clone ID: LIB3061-002-Q1-K1-D12
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359 gttatattcg ccaagactgg gctggattac gtttcttgct cccctttcag ggtcccgatc   120
361 gctaggctag ctgcagctca ggtgcttgct tgaggggtgcc tctgctcgtt ggcagccgga   180
363 tcgcctgctg ttggtgcacg ttgtgattaa taatactact atgacagagc catgctctgt   240
365 gaagagtatt agtagcagcg ctcataaaag ctacagttcc gtctctattc tcagctatgt   300
367 aaaacttcca aactgttcat gcttaaaact gaagggtttc ttggtgtgag gtgtgcatgt   360
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372 <210> SEQ ID NO: 16
373 <211> LENGTH: 419
374 <212> TYPE: DNA
375 <213> ORGANISM: Zea mays
377 <223> OTHER INFORMATION: Clone ID: LIB3061-049-Q1-K1-C8
379 <400> SEQUENCE: 16
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383 tcgggagccc cgacgatggc caccaccgcg gtgtcggtcg acgagaagct cgataagctt   120
385 cgcgcgaggg tcgccaagct cgaccagatc agcgagaacg agaagtccgg gttcatcagc   180
387 ctctgtgtcac ggtacctcag tggggaggcg gagcagatcg agtggagcaa gatccagacc   240
389 cctacggatg aagtgggtgt gcctacgat accgtcgcgt cgctctccga agatctcgag   300
391 gagacgaaga agctgctgga taagctcgtt gtgtcgaagc ttaacggagg gctcgggacg   360
393 accatgggct gcaactgggc caagtctgtc attgaagtcc gcaatgggtt cacattcct   419
396 <210> SEQ ID NO: 17
397 <211> LENGTH: 431
398 <212> TYPE: DNA
399 <213> ORGANISM: Zea mays
401 <223> OTHER INFORMATION: Clone ID: LIB3061-026-Q1-K1-D7
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405 ctggcagtgta ctattcgact tctggaccct cccctccacg agttccttcc agaagggggc   60
407 gtcgaggaaa tcgtgcgtga actgtgttct gagacgggag ccaaccagga ggatgccctc   120
409 gcacggatcg aaaagctttc ggaagtaaac ccgatgcttg gcttcctgtg ctgcaggctt   180
411 ggtatatcgt accctgagct aacagaaatg caagcccgtg ccatctttga agctgtata   240
413 gcgatgacca accagggcgt tcaagtcttt ccagagataa tggttcctct tgttgaaca   300
415 ccgcaggaat tgggacatca ggtggccctt atccgtcaag tcgctaacaa agttttcacc   360
417 agtatgggga aaactattgg gtacaagatc ggaacgatga ttgagattcc cagggcagct   420
419 ctagtggctg a
422 <210> SEQ ID NO: 18
423 <211> LENGTH: 319
424 <212> TYPE: DNA
425 <213> ORGANISM: Zea mays
427 <223> OTHER INFORMATION: Clone ID: LIB3061-009-Q1-K1-F12
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433 gggttgtgcc ggacagagca catgttcttc gcttcggatg agaggattaa ggctgtgagg   120
435 cagatgatta tggctcccac agttgagctg aggcagcaag cgctggatcg ccttctgcct   180
437 taccagaggt ctgacttcga aggcattctc cgtgctatgg atggactttc agtgactatt   240
439 cgacttctgg accctccact acacgagttc ctttcagaag ggaacgtcga ggaaatcgtg   300

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**Please Note:**

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

## VERIFICATION SUMMARY

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Input Set : D:\AprilCornReg.rpt

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L:9 M:270 C: Current Application Number differs, Replaced Current Application No  
L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
L:199 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:8  
L:199 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:8  
L:199 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:8  
L:199 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:8  
L:212 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:9  
L:212 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:9  
L:212 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:9  
L:212 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:9  
L:341 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:14  
L:341 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:14  
L:341 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:14  
L:341 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:14  
L:488 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:20  
L:488 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:20  
L:488 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:20  
L:488 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:20  
L:861 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:36  
L:861 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:36  
L:861 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:36  
L:861 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:36  
L:1090 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:46  
L:1090 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:46  
L:1090 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:46  
L:1090 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:46  
L:1143 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:48  
L:1143 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:48  
L:1143 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:48  
L:1143 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:48  
L:1290 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:54  
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L:1290 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:54  
L:1290 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:54  
L:1405 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:59  
L:1405 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:59  
L:1405 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:59  
L:1405 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:59  
L:1422 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:60  
L:1422 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:60  
L:1422 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:60  
L:1422 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:60  
L:1455 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:61  
L:1455 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:61  
L:1455 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:61  
L:1455 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:61  
L:1578 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:66  
L:1578 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:66

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L:1578 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:66  
L:1578 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:66  
L:1629 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:68  
L:1629 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:68  
L:1629 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:68  
L:1629 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:68  
L:1654 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:69  
L:1654 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:69  
L:1654 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:69  
L:1654 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:69  
L:1656 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:69  
L:1656 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:69  
L:1656 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:69  
M:340 Repeated in SeqNo=69  
L:1683 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:70  
L:1683 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:70  
L:1683 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:70  
L:1683 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:70  
L:1776 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:74  
L:1776 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:74  
L:1776 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:74  
L:1919 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:80  
M:340 Repeated in SeqNo=80  
L:2010 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:84  
L:2065 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:86  
M:340 Repeated in SeqNo=86  
L:2808 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:116  
L:2895 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:120  
L:2940 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:122  
L:3239 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:135  
L:3598 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:150  
M:340 Repeated in SeqNo=150  
L:4291 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:179  
L:5294 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:219  
L:5403 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:224  
L:5428 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:225  
L:5649 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:234  
L:5700 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:236  
L:5725 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:237  
L:5796 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:240  
L:5813 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:241  
L:6274 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:260  
L:6343 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:263  
L:6370 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:264  
L:6539 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:271  
L:6568 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:272  
L:6615 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:274  
L:6672 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:277  
L:6749 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:280

VERIFICATION SUMMARY

DATE: 05/09/2000

PATENT APPLICATION: US/09/553,094

TIME: 16:28:07

Input Set : D:\AprilCornReg.rpt

Output Set: N:\CRF3\05092000\I553094.raw

L:6850 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:284  
M:340 Repeated in SeqNo=284  
L:6875 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:285  
L:7246 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:300  
L:7275 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:301